

ABSTRACT

A method for measuring a target nucleic acid, which does not require skill and which enables to detect a SNP and to quantify the target nucleic acid simply in a short time is disclosed. In the method of the present invention, the target nucleic acid, a labeled probe which is a labeled nucleic acid which hybridizes with the target nucleic acid, a non-labeled probe which is a nucleic acid having a nucleotide sequence complementary to a region in the target nucleic acid, which region is different from the region with which the labeled probe hybridizes, and an immobilized probe which is a nucleic acid bound to a support, which nucleic acid has a nucleotide sequence complementary to a region in the target nucleic acid, which region is different from the region with which the labeled probe hybridizes are reacted; and the label of the labeled probe bound to the support is measured. An end region of the region with which the non-labeled probe hybridizes overlaps with an end region of the region with which the immobilized probe hybridizes.